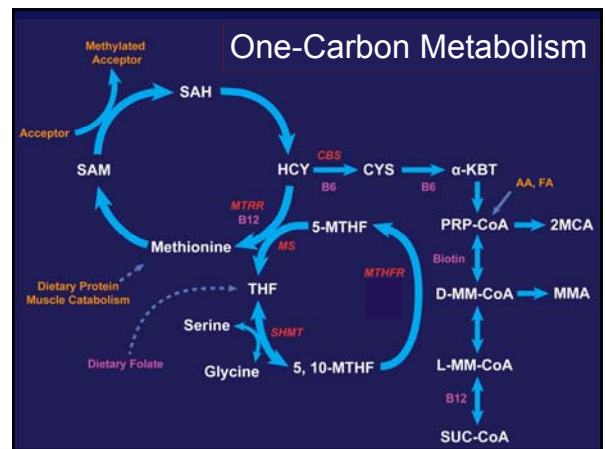
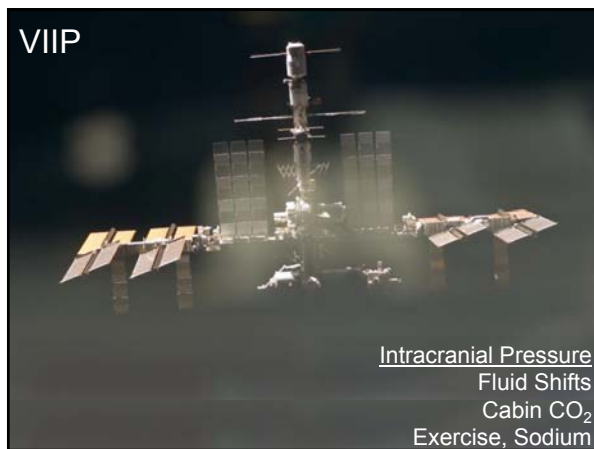
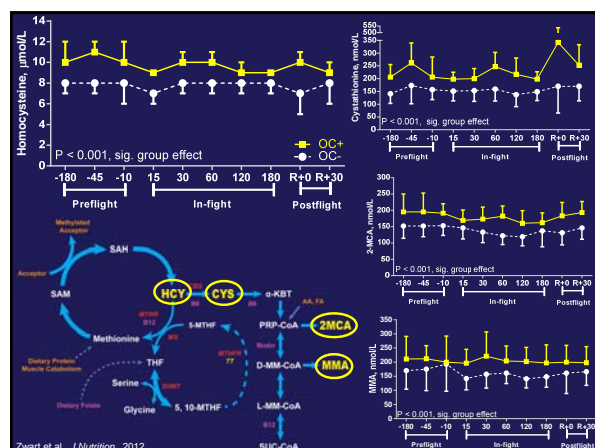


### Forward Work

**Bone Strength?**  
**Fracture risk?**

**Optimization**  
**Exercise**  
**Diet**





# MTHFR C677T Polymorphism

The diagram illustrates the MTHFR C677T polymorphism and its impact on the methionine cycle. On the left, two DNA double helices are shown. The top helix represents the wild-type (C/C) genotype, with green 'C' alleles at the 677th position. The bottom helix represents the mutant (T/T) genotype, with yellow 'T' alleles at the 677th position. In the center, a table summarizes the prevalence and enzymatic activity of these genotypes. On the right, a metabolic pathway diagram shows the conversion of SAM to SAH, which is then recycled back to SAM by Methylenetetrahydrofolate Reductase (MTHFR). The MTHFR enzyme is shown in two states: a green state for the wild-type (C/C) and a yellow state for the mutant (T/T). The mutant enzyme is shown with a red 'X' over it, indicating reduced activity. The pathway continues from SAM to Methionine, which is then converted to S-adenosylmethionine (SAM) by Methionine Adenosyltransferase (MAT). SAM is then used for various methylation reactions, including the methylation of DNA, RNA, and proteins. The diagram also shows the conversion of SAM to S-adenosylhomocysteine (SAH) by Methyltransferase (MT), which is then converted back to Homocysteine by S-adenosylhomocysteine Lyase (AHCY). Homocysteine is then converted to Methionine by Methionine Synthase (MS), which uses 5,10-methylenetetrahydrofolate (5,10-MTHF) as a cofactor. The diagram also shows the conversion of Homocysteine to Methionine by Methionine Synthase (MS), which uses 5,10-methylenetetrahydrofolate (5,10-MTHF) as a cofactor. The diagram also shows the conversion of Homocysteine to Methionine by Methionine Synthase (MS), which uses 5,10-methylenetetrahydrofolate (5,10-MTHF) as a cofactor.

	% Pop	Enz Act.
C/C	~35%	100%
C/T(T/C)	~50%	~66%
T/T	~15%	~50%

The diagram also shows the conversion of SAM to SAH by Methyltransferase (MT), which is then converted back to SAM by Methylenetetrahydrofolate Reductase (MTHFR). The MTHFR enzyme is shown in two states: a green state for the wild-type (C/C) and a yellow state for the mutant (T/T). The mutant enzyme is shown with a red 'X' over it, indicating reduced activity. The pathway continues from SAM to Methionine, which is then converted to S-adenosylmethionine (SAM) by Methionine Adenosyltransferase (MAT). SAM is then used for various methylation reactions, including the methylation of DNA, RNA, and proteins. The diagram also shows the conversion of SAM to S-adenosylhomocysteine (SAH) by Methyltransferase (MT), which is then converted back to Homocysteine by S-adenosylhomocysteine Lyase (AHCY). Homocysteine is then converted to Methionine by Methionine Synthase (MS), which uses 5,10-methylenetetrahydrofolate (5,10-MTHF) as a cofactor. The diagram also shows the conversion of Homocysteine to Methionine by Methionine Synthase (MS), which uses 5,10-methylenetetrahydrofolate (5,10-MTHF) as a cofactor. The diagram also shows the conversion of Homocysteine to Methionine by Methionine Synthase (MS), which uses 5,10-methylenetetrahydrofolate (5,10-MTHF) as a cofactor.

# MTRR A66G

The diagram illustrates the metabolic pathways involving folate and methionine, with a focus on the MTRR A66G mutation. Key components include:

- Methionine Cycle:** SAM (S-adenosylmethionine) is converted to SAH (S-adenosylhomocysteine) by a methyltransferase (Methylated Acceptor). SAH is then converted back to SAM by MTRR (Methylenetetrahydrofolate reductase), which uses 5-MTHF (5-methyltetrahydrofolate) as a cofactor. This step is highlighted with a yellow oval and labeled "MTRR B12".
- Transsulfuration Pathway:** HCY (homocysteine) is converted to CYS (cysteine) by CBS (cystathionine beta-synthase), which uses B6 (pyridoxal phosphate) as a cofactor. CYS is then converted back to HCY by α-KBT (α-ketobutyrate transaminase), which uses B6 and AA (ascorbic acid) as cofactors.
- Folate Pathway:** THF (tetrahydrofolate) is converted to 5-MTHF by MTRR. 5-MTHF is then converted to 5,10-MTHF by MS (methylenesuccinyl-CoA synthetase). 5,10-MTHF is converted back to 5-MTHF by MTHFR (methylenetetrahydrofolate reductase), which uses TT (thiamine) as a cofactor. A dashed line indicates that MTHFR is also involved in the conversion of 5-MTHF to THF.
- Other Pathways:** PRP-CoA (phosphoribosyl-CoA) is converted to 2MCA (2-methylcrotonyl-CoA) by Biotin. D-MM-CoA (dimethylmalonyl-CoA) is converted to MMA (methylmalonyl-CoA). L-MM-CoA (l-methylmalonyl-CoA) is converted to SUC-CoA (succinyl-CoA) by B12 (cobalamin).
- Regulation:** Dietary Protein and Muscle Catabolism are shown to influence the folate pathway. Dietary Folate is shown to influence the conversion of THF to 5-MTHF.

